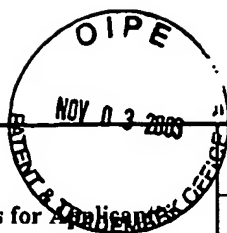


Form PTO-1449 (modified)

List of Patents and Publications for Applicant

INFORMATION DISCLOSURE STATEMENT

(Use several sheets if necessary)

Atty. Docket No.
AMBI:063USSerial No.
10/632,539Applicant
Matthew M. Winkler and David BrownFiling Date:
July 31, 2003Group:
UnknownU.S. Patent Documents
See Page 1Foreign Patent Documents
See Page 1Other Art
See Page 1

U.S. Patent Documents

Exam. Init.	Ref. Des.	Document Number	Date	Name	Class	Sub Class	Filing Date of App.
SC	A1	5,262,311	11/16/93	Pardee <i>et al.</i>	435	91.2	03/11/92
SC	A2	5,545,522	08/13/96	Van Gelder <i>et al.</i>	435	6	10/05/92
SC	A3	5,861,245	01/19/99	McClelland	435	6	06/06/95
SC	A4	6,132,997	10/17/200	Shannon	435	91.21	05/28/99

Foreign Patent Documents

Exam. Init.	Ref. Des.	Document Number	Date	Country	Class	Sub Class	Translation Yes/No
SC	B1	EP 0 416 817	03/13/91	Europe			
SC	B2	EP 0 870 842	10/14/98	Eurpoe			
SC	B3	WO 00/05409	02/03/00	PCT			
SC	B4	WO 00/24939	05/04/00	PCT			
SC	B5	WO 00/75356	12/14/00	PCT			
SC	B6	WO 02/064835	01/31/02	PCT			
SC	B7	WO 97/27317	07/31/97	PCT			
SC	B8	WO 98/08973	03/05/98	PCT			

Other Art (Including Author, Title, Date Pertinent Pages, Etc.)

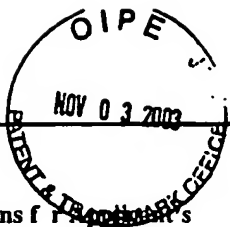
Exam. Init.	Ref. Des.	Citation
SC	C1	Kato, "Adaptor-tagged competitive PCR: a novel method for measuring relative gene expression," <i>Nucleic Acids Research</i> , Oxford University Press, Surrey, GB, 25(22):4694-4696, 1997.
SC	C2	Kita <i>et al.</i> , "Modulation of polyglutamine-induced cell death by genes identified by expression profiling," <i>Human Molecular Genetics</i> , 11(19):2279-2287, 2002.

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Other Art (Including Author, Title, Date Pertinent Pages, Etc.)

Exam. Init.	Ref. Des.	Citation
SC	C3	Matoba <i>et al.</i> , "Gene expression in mouse cerebellum during its development," <i>Gene</i> , 241:125-131, 2000.
SC	C4	Matoba <i>et al.</i> , "Gene expression profiling of mouse postnatal cerebellar development," <i>Physiol. Genomics</i> , 4:155-164, 2000.
SC	C5	Phillips <i>et al.</i> , "Antisense RNA amplification: A linear amplification method for analyzing the mRNA population," <i>Methods, a Companion to Methods in Enzymology</i> , 10(3):283-288, 1996.
SC	C6	Sakai <i>et al.</i> , "Microarray hybridization with fractionated cDNA: enhanced identification of differentially expressed genes," <i>Analytical Biochemistry</i> , 287(1):32-37, 2000.
SC	C7	Shuldiner <i>et al.</i> , "RNA template-specific polymerase chain reaction RS-PCR a novel strategy to reduce dramatically false positives," <i>Gene</i> , 91(1):139-142, 1990.
SC	C8	Welch <i>et al.</i> , "Fingerprinting genomes using PCR with arbitrary primers," <i>Nucleic Acids Research</i> , Oxford University Press, Surrey, GB, 18(24):7213-7218, 1990.
SC	C9	Welch <i>et al.</i> , "Nucleic acid fingerprinting by PCR-based methods: applications to problems in aging and mutagenesis," <i>Mutation Research</i> , 338(1-6):215-229, 1995.
SC	C10	Wyttenbach <i>et al.</i> , "Polyglutamine expansions cause decreased CRE-mediated transcription and early gene expression changes prior to cell death in an inducible cell model of Huntington's disease," <i>Human Molecular Genetics</i> , 10(17):1829-1845, 2001.
SC	C11	Zimmerman <i>et al.</i> , "Technical aspects of quantitative competitive PCR," <i>Biotechniques</i> , 21(2):268-270, 1996.

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